

03610365-031604

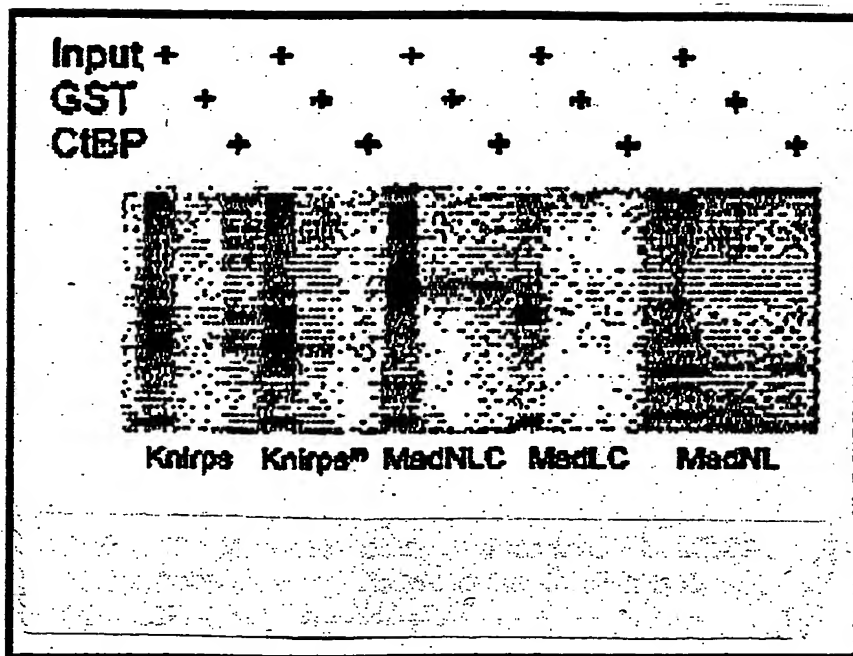


FIGURE 1

Smad4	20	50	100	100	100
dCtBP	-	-	-	50	100

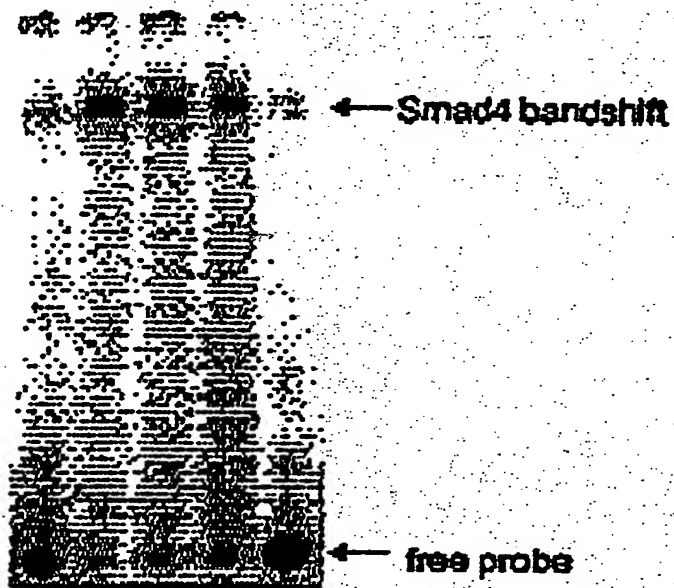


FIGURE 2

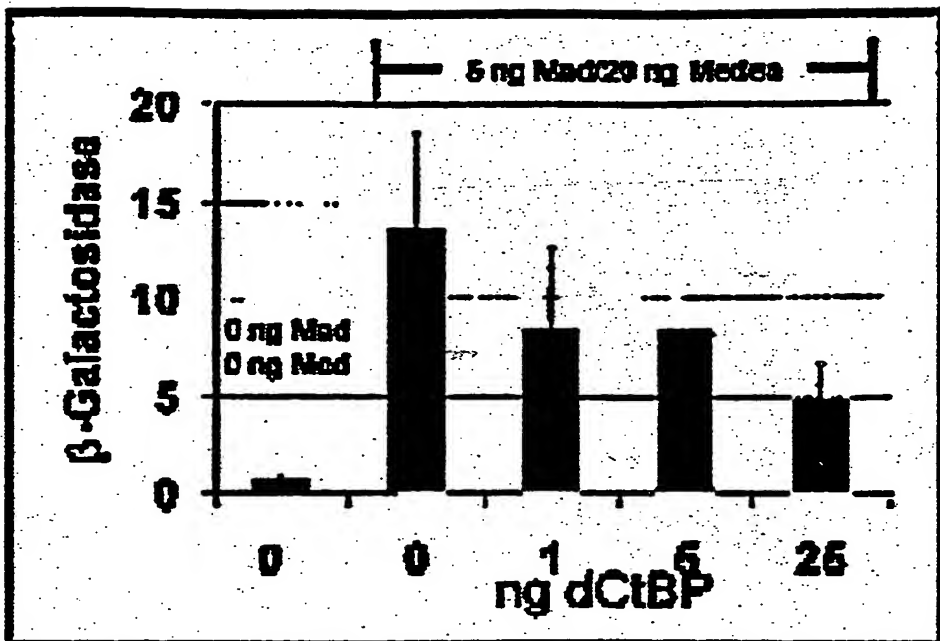


FIGURE 3

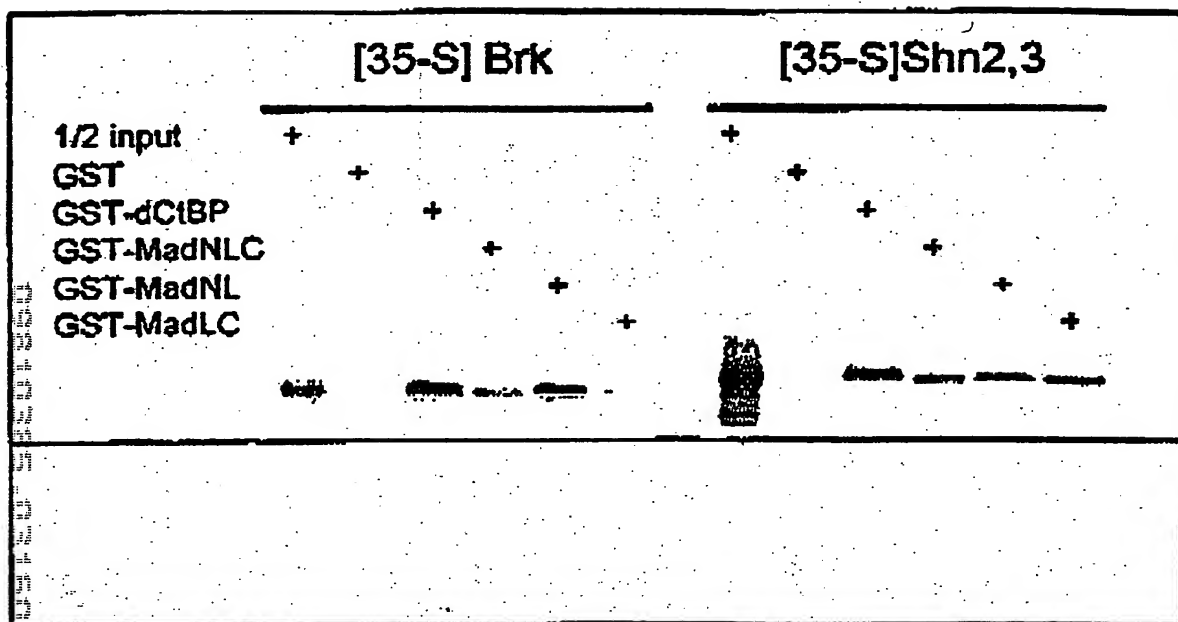


FIGURE 4

006102305-02160-1

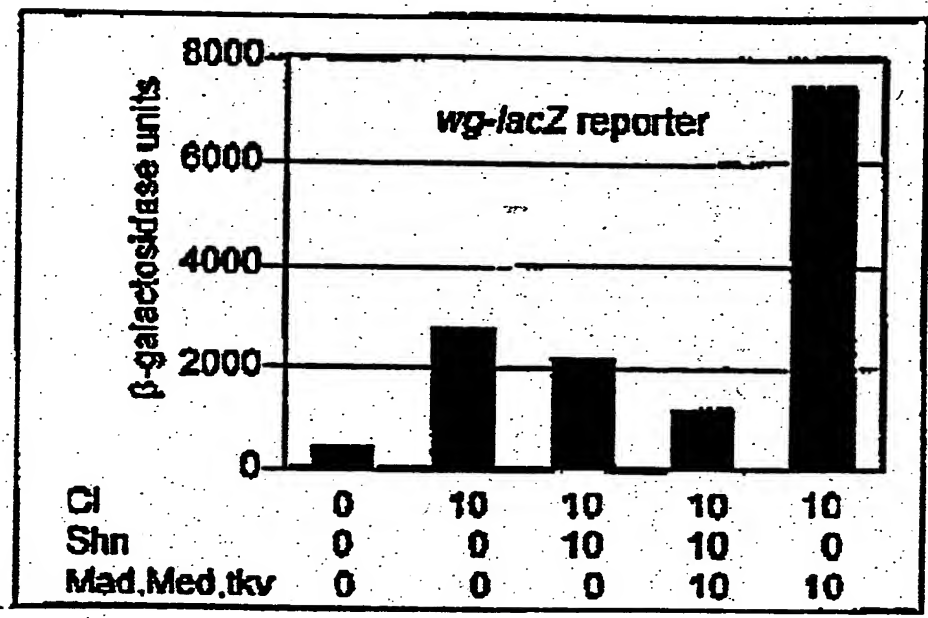


FIGURE 5

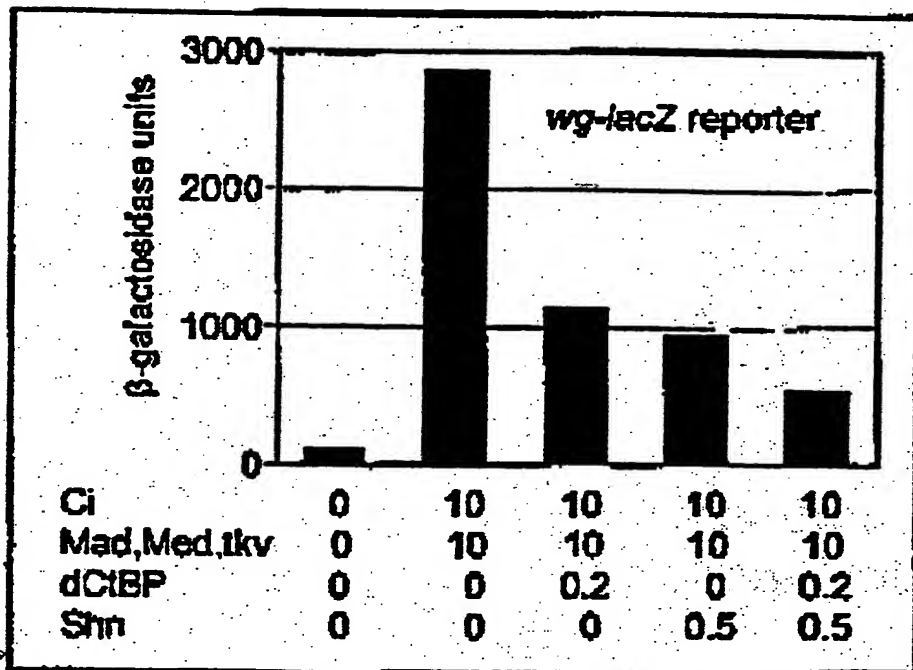


FIGURE 6

Schematic representation of the organization of the genes Bklf, Evi-1, AREB5, Zfh-1, FOG, Ush, Kr, Shn, and Brk. Each gene is represented by a horizontal line with vertical bars indicating exons and circles with crosses indicating introns. A scale bar at the bottom indicates 500 bp.

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

DRQP	QVE	FVDLT	VNKR
Evi-1	SEB	PFDLT	TKRK
AREB6	QDQ	PLDLS	MGSR
SIF1	QSE	PLNLS	CAKK
	TPS	PLNLS	STSG
	QAE	FLDLS	LPKO
	SDS	PLNLT	VTKK
ZEB	SPS	PLNLS	SSRN
	QVE	PLDLS	LKQQ
FOG	PDG	PIDLS	KPRR
CtIP	MDK	PLDLS	DRFS
Kruppal	EEA	PLDLS	EDGA
	QTE	MDLS	MHSF
Zeh-1	EDQ	PLDLS	VKRD
Ush	OEG	MDLS	MHST
Shn	SKN	PKQLS	KSKS
	PVM	PLNLS	AKPK
	ESA	MDLT	KPRG
Brk	TPA	MDLS	GSEA
	DTA	PTNLT	LVA
Kajrps	QEG	MDLS	MKTS
Snail	QDQ	PDLS	LKRG
	CSE	FEDLS	VRND
Hairy	EQO	PLSLV	TKQQ
E(spl)m5	APZ	FVNLA	DOXR
Ad2 243R	PGQ	PLDLS	CKRP
Ad12 235R	QTV	FVDLS	VKRP
TGIF	MDI	PLDLS	SSAG
	PFT	PFDLN	QDL'S
Consensus	tee	PIDLS	Xkkk
	dc	vn t	rrr

FIGURE 8